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OM protein - protein search, using sw model

Run on: July 16, 2003, 13:53:06 ; Search time 69 Seconds
(Without alignments)
42.486 Million cell updates/sec

Title: US-09-914-213-2

Perfect score: 116
Sequence: 1 GLEISEINEEDLKECFDDME 22

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 908470 seqs, 133250620 residues

Total number of hits satisfying chosen parameters: 908470

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%

Listing first 45 summaries

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22: /SID2/gcgdata/geneseq/geneseq-emb1/AA2001.DAT:*
23: /SID2/gcgdata/geneseq/geneseq-emb1/AA2002.DAT:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length DB	ID	Description
1	116	100.0	22	AA822023	Human CFTR peptide
2	116	100.0	22	AA831723	A human cystic fib
3	116	100.0	22	AB617640	Novel human diago
4	116	100.0	1843	ABG14961	Novel human diago
5	110	94.8	1091	AA13303	CFTR Y1092X. Homo
6	110	94.8	1190	AA13308	CFTR 3659 del C.
7	110	94.8	1476	AA13368	CFTR protein seque
8	110	94.8	1479	AA11602	Mutant cystic fibr
9	110	94.8	1479	AA13321	CFTR delta 1507.
10	110	94.8	1479	AA102279	DeltaF508 cystic f

11	110	94.8	1479	AA74516	Human delta-F508-C
12	110	94.8	1480	AA11115	Cystic fibrosis tr
13	110	94.8	1480	AA13252	CFTR G85E. Homo s
14	110	94.8	1480	AA13253	CFTR I148T. Homo
15	110	94.8	1480	AA13254	CFTR G178R. Homo
16	110	94.8	1480	AA13255	CFTR A455E. Homo
17	110	94.8	1480	AA13297	CFTR S549R. Homo
18	110	94.8	1480	AA13298	CFTR G551D. Homo
19	110	94.8	1480	AA13299	CFTR R560T. Homo
20	110	94.8	1480	AA13300	CFTR Y574H. Homo
21	110	94.8	1480	AA13302	CFTR L1077P. Homo
22	110	94.8	1480	AA13302	Cystic fibrosis tr
23	110	94.8	1480	AA13302	Cystic fibrosis tr
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43	110	94.8	1480	AA13302	Cystic fibrosis tr
44	110	94.8	1480	AA13302	Cystic fibrosis tr
45	110	94.8	1480	AA13302	Cystic fibrosis tr

ALIGNMENTS

RESULT 1
AA822023 standard; peptide; 22 AA.
XX
AC AA822023:
XX
DT 08-JAN-2001 (first entry)
XX
DE Human CFTR peptide fragment NEG2.
XX
KW Human; cystic fibrosis transmembrane conductance regulator; CFTR; CF;
KW chloride channel.
XX
OS Homo sapiens.
XX
PN WO200050591-A1.
XX
XX 31-AUG-2000.
XX
PD 24-FEB-2000; 2000WO-US04642.
XX
PF 24-FEB-1999; 99US-0121495.
XX
PR (UYCA-) UNIV CASE WESTERN RESERVE.
XX
PA Adams LM, Davis PB, Ma J;
XX
PI WPI: 2000-572090/53.
XX
DR Polypeptide useful for enhancing the open probability of cystic
XX fibrosis transmembrane conductance regulator chloride channel in cystic
XX fibrosis patients having minimally active mutant protein

PS Claim 2; Fig 2; 35pp; English.
 CC Defects in the cystic fibrosis transmembrane conductance regulator
 CC (CFTR), are associated with cystic fibrosis (CF). CFTR is a chloride
 CC channel located in the apical membrane of epithelial cells. The present
 CC peptide is a fragment of human CFTR. This peptide is useful for
 CC activating and opening a CFTR protein by the formation of cAMP regulated
 CC chloride channel. This peptide can therefore be used as therapy for CF.
 XX
 SQ Sequence 22 AA;
 Query Match 100.0%; Score 116; DB 21; Length 22;
 Best Local Similarity 100.0%; Pred. No. 2e-09;
 Matches 22; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 1 GLEISEEINEDLKCEFFDME 22
 DB 1 GLEISEEINEDLKCEFFDME 22
 RESULT 2
 AAB31723 standard; Protein; 1480 AA.
 AC AAB31723;
 XX 30-APR-2001 (first entry)
 DT
 DE A human cystic fibrosis transmembrane conductance regulator.
 XX
 XX Arginine-framed tripeptide; AFT; endoplasmic reticulum retention;
 KW Cystic fibrosis transmembrane conductance regulator; CFTR; hormone;
 KW Cystic fibrosis; macular dystrophy; stargardt's disease; growth factor;
 KW export-incompetent protein; adenine-nucleotide binding cassette protein;
 KW ABC protein; immune regulator; adhesion protein; clotting factor;
 KW hemostatic regulator; intracellular transport.
 KM
 XX Homo sapiens.
 OS
 XX WO200103722-A1.
 PN 18-JAN-2001.
 PD
 XX 07-JUL-2000; 2000WO-US40324.
 PF
 XX 09-JUL-1999; 99US-0143090.
 PR 21-JUL-1999; 99US-0146097.
 XX
 XX (MAYO-) MAYO FOUND MEDICAL EDUCATION & RES.
 PA
 XX
 PI Rlordan JR, Chang X;
 DR WPI; 2001-123141/13.
 XX
 PT Novel cystic fibrosis transmembrane conductance regulator proteins,
 PT useful for treating cystic fibrosis have one arginine, in an arginine
 PT tripeptide sequence motif, substituted by other amino acid
 PS Disclosure; Fig 6; 56pp; English.
 XX
 XX The present sequence represents a human cystic fibrosis transmembrane
 CC conductance regulator (CFTR). Arginine-framed tripeptide (AFT) sequences
 CC of CFTR contribute to endoplasmic reticulum (ER) retention or delay in
 CC maturation of proteins. The AFT sequences are useful as competitive
 CC inhibitors of ER retention. The peptides are useful for treating cystic
 CC fibrosis. They are useful for treating a subject having a suspected of
 CC having a physiological disorder (e.g. macular dystrophy and stargardt's
 CC disease) associated with an export-incompetent protein such as
 CC adenine-nucleotide binding cassette (ABC) protein, growth factor, immune
 CC regulator, adhesion protein, hormone, clotting factor, hemostatic
 CC regulator and their receptors. The AFT peptides are also useful for
 CC inducing or increasing intracellular transport of an export-incompetent
 CC protein in a cell e.g. a cell surface or secreted protein and preferably

CC export-incompetent CFTR. They are also useful for inhibiting degradation
 CC of a secreted or cell surface protein in a cell and for detecting the
 CC presence of an export-incompetent protein in a cell.
 XX
 SQ Sequence 1480 AA;
 Query Match 100.0%; Score 116; DB 22; Length 1480;
 Best Local Similarity 100.0%; Pred. No. 1.7e-07;
 Matches 22; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 QY 1 GLEISEEINEDLKCEFFDME 22
 DB 817 GLEISEEINEDLKCEFFDME 838
 RESULT 3
 ABG17640 standard; Protein; 1614 AA.
 ID ABG17640
 XX
 XX ABG17640;
 AC
 XX 18-FEB-2002 (first entry)
 DT
 DE Novel human diagnostic protein #17631.
 XX
 XX Human; chromosome mapping; gene mapping; gene therapy; forensic;
 KW food supplement; medical imaging; diagnostic; genetic disorder.
 KM
 XX Homo sapiens.
 OS
 XX WO200175067-A2.
 PN 11-OCT-2001.
 PD
 XX 30-MAR-2001; 2001WO-US08631.
 PF
 XX 31-MAR-2000; 2000US-0540217.
 PR 23-AUG-2000; 2000US-0649167.
 XX
 XX (HYSE-) HYSEQ INC.
 PA
 XX Drmanac RT, Liu C, Tang YT;
 PI WPI; 2001-639362/73.
 DR N-PSDB; AAS81827.
 XX
 XX New isolated polynucleotide and encoded polypeptides, useful in
 PT diagnostics, forensics, gene mapping, identification of mutations
 PT responsible for genetic disorders or other traits and to assess
 PT biodiversity
 PS
 XX Claim 20; SEQ ID No 47999; 103pp; English.
 XX
 XX The invention relates to isolated polynucleotide (I) and
 CC polypeptide (II) sequences. (I) is useful as hybridisation probes,
 CC polymerase chain reaction (PCR) primers, oligomers, and for chromosome
 CC and gene mapping, and in recombinant production of (II). The
 CC polynucleotides are also used in diagnostics as expressed sequence tags
 CC for identifying expressed genes. (I) is useful in gene therapy techniques
 CC to restore normal activity of (II) or to treat disease states involving
 CC (II). (II) is useful for generating antibodies against it, detecting or
 CC quantitating a polypeptide in tissue, as molecular weight markers and as
 CC a food supplement. (II) and its binding partners are useful in medical
 CC imaging of sites expressing (II). (I) and (II) are useful for treating
 CC disorders involving aberrant protein expression or biological activity.
 CC The polypeptide and polynucleotide sequences have applications in
 CC diagnostics, forensics, gene mapping, identification of mutations
 CC responsible for genetic disorders or other traits to assess biodiversity
 CC and to produce other types of data and products dependent on DNA and
 CC amino acid sequences. ABG00010-ABG30377 represent novel human
 CC diagnostic amino acid sequences of the invention.
 CC Note: The sequence data for this patent did not appear in the printed
 CC specification, but was obtained in electronic format directly from WIPO


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FT 582 Modified-site /label= phosphorylation_site /note= "by protein kinases C"
FT 623 Modified-site /label= phosphorylation_site /note= "by protein kinases C"
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FT 707 Modified-site /label= phosphorylation_site /note= "by protein kinases C"
FT 712 Modified-site /label= phosphorylation_site /note= "by protein kinases A"
FT 737 Modified-site /label= phosphorylation_site /note= "by protein kinases A"
FT 768 Modified-site /label= phosphorylation_site /note= "by protein kinases C"
FT 788 Modified-site /label= phosphorylation_site /note= "by protein kinases A"
FT 790 Modified-site /label= phosphorylation_site /note= "by protein kinases C"
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FT WO9110734-A.
FT 25-JUL-1991.
FT 11-JAN-1991. 91WO-CA00009.
FT 10-JUL-1990. 90CA-2020817.
FT 12-JAN-1990. 90CA-2007699.
FT 01-MAR-1990. 90CA-2011253.
XX (HSCR-) HSC RES DEV CORP.
XX Tsui LC, Rommens JM, Kerem B:
XX WPT; 1991-238022/32.
XX N-PDB; AAQ13066.
XX DR
XX PT Mutant cystic fibrosis trans-membrane conductance regulator gene
XX of cystic fibrosis
XX PS Claim 20; Page 124; 178pp; English.
XX CC In the LY1092X mutation a C to A change is detected at nucleotide
XX position 3408. This would result in protein synthesis termination
XX at amino position 1092. Hence the amino acid Tyr is not present
XX CC In the truncated polypeptide.
XX CC The mutant CF gene when expressed in cells of the human body, is
XX associated with altered cell function which correlates with the
XX genetic disease cystic fibrosis.
XX See also AAQ13053-72.
XX
XX SQ Sequence 1091 AA:
XX
XX Query Match 94.8%; Score 110; DB 12; Length 1091;
XX Best Local Similarity 95.5%; Pred. No. 8.2e-07;
XX Matches 21; Conservative 0; Mismatches 1; Indels 0; Gaps 0;
QY 1 GLEISEINEDIKCFDDME 22
DB 817 GLEISEINEDIKCFDDME 838
RESULT 6
AAR13308 standard; Protein; 1190 AA.
XX AAR13308;
XX 14-OCT-1991 (first entry)

```

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DE		Deletion; mutant; diagnosis; antibodies; drug therapy.	
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KW		Homo sapiens.	
OS			
XX			
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XX	25-JUL-1991.
PD	
XX	
PE	11-JAN-1991; 91WO-C400009.
PR	10-JUL-1990; 90CA-2020817.
PR	12-JAN-1990; 90CA-2007659.
PR	01-MAR-1990; 90CA-2011253.
XX	
PA	(HSCR-) HSC RES DEV CORP.
XX	
PI	Tsuji LC, Rommens JM, Kerem B;
DR	WPI: 1991-238022/32.
DR	N-PSDB; AAQ13072.
XX	
PT	Mutant cystic fibrosis trans-membrane conductance regulator gene
PT	-used for producing prods. for diagnosis, screening and therapy
PT	of cystic fibrosis
SS	Claim 20: Page 124; 178op: English).

Claim 20; Page 124; 178pp; English.

XX 3659 del C is a frameshift mutation in exon 19.
 CC The 3659 del C mutation results in a shortened polypeptide
 CC significantly different from the single amino acid deletions or
 CC alterations.
 CC The mutant CF gene when expressed in cells of the human body, is
 CC associated with altered cell function which correlates with the
 CC genetic disease cystic fibrosis.
 CC See also AAQ13053-72.

XX Sequence 1190 AA:

Query Match 94.8%; Score 110; DB 12; Length 1190;

Best Local Similarity 95.5%; Pred. No. 9e-07; Mismatches 0; Gaps 0;

DB 817 GLEISEINEEDLKECFDDME 22
 1 GLEISEINEEDLKECFDDME 22
 817 GLEISEINEEDLKECFDDME 838

RESULT 7

AAV33968 standard; Protein; 1476 AA.

AAV33968;

26-MAY-2000 (first entry)

CFTR protein sequence.

AAV vector; inverted terminal repeat; ITR; gene therapy; CFTR; TK gene;
 cystic fibrosis transmembrane conductance regulator; cystic fibrosis;
 promoter; HSV; thymidine kinase; chromosome 7q31.

Homo sapiens.

Key Location/Qualifiers

Misc-difference 1150 /note="encoded by AGC"

WO9943789-A1.

02-SEP-1999.

25-FEB-1999; 99WO-US04212.

25-FEB-1998; 98US-0075980.

(REGC) UNIV CALIFORNIA.

Dong J, Kan YW;

WPI; 1999-550866/46.

N-PSDB; AAZ11643.

Efficient AAV vectors useful in gene therapy protocols for the

treatment of cystic fibrosis

Example 1; Page 33; 34pp; English.

The invention provides efficient AAV vectors with improved capacity for
 DNA due to the removal of all nucleic acid sequences that are not
 essential for replication (to leave just 2 inverted terminal repeat
 sequences (ITRs)). The AAV vectors may be used for the delivery of
 therapeutic nucleic acids in gene therapy protocols. In particular,
 they may be used to deliver cystic fibrosis (CF) transmembrane
 conductance regulator (CFTR) polynucleotides to the respiratory tract
 of CF patients to rectify mutations in the patients own CFTR genes and
 restore normal function to the chloride channel for replication,
 vector lacks all nucleic acids that are not essential for replication,
 therefore giving it a greater capacity for exogenous DNA and hence
 improving the efficiency with which it transfects cells. The AAV vectors

CC of the invention can efficiently and persistently transfer CFTR
 CC polynucleotides to the airway epithelium of CF patients without any
 CC adverse side effects. The present sequence represents the CFTR protein.
 XX

SO Sequence 1476 AA:

Query Match 94.8%; Score 110; DB 20; Length 1476;

Best Local Similarity 95.5%; Pred. No. 1.1e-06; Mismatches 0; Gaps 0;

DB 817 GLEISEINEEDLKECFDDME 22
 1 GLEISEINEEDLKECFDDME 22
 817 GLEISEINEEDLKECFDDME 838

RESULT 8

AA11602 standard; Protein; 1479 AA.

AA11602;

22-MAY-1991 (first entry)

Mutant cystic fibrosis transmembrane conductance regulatory protein.
 Cystic fibrosis; transmembrane conductance regulatory protein; CFTR;
 diagnosis; mutant.

Homo sapiens.

Key Location/Qualifiers

Misc-difference 81..102 /label="potential_membrane-spanning_segment

Misc-difference 118..138 /label="potential_membrane-spanning_segment

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FT	Misc-difference	681..681	/label- protein_kinase_C-phosphorylation_site
FT	Misc-difference	685..685	/label- protein_kinase_C-phosphorylation_site
FT	Misc-difference	706..706	/label- protein_kinase_C-phosphorylation_site
FT	Misc-difference	779..779	/label- protein_kinase_C-phosphorylation_site
FT	Misc-difference	790..790	/label- protein_kinase_C-phosphorylation_site
FT	Misc-difference	808..808	/label- protein_kinase_C-phosphorylation_site
FT	Misc-difference	965..965	/label- protein_kinase_C-phosphorylation_site
FT	Misc-difference	1057..1057	/label- protein_kinase_C-phosphorylation_site
FT	Misc-difference	1063..1063	/label- protein_kinase_C-phosphorylation_site
FT	Misc-difference	1094..1094	/label- protein_kinase_C-phosphorylation_site
FT	Misc-difference	1177..1177	/label- protein_kinase_C-phosphorylation_site
FT	Misc-difference	1210..1210	/label- protein_kinase_C-phosphorylation_site
FT	Misc-difference	1215..1215	/label- protein_kinase_C-phosphorylation_site
FT	Misc-difference	1247..1247	/label- protein_kinase_C-phosphorylation_site
FT	Misc-difference	1298..1298	/label- protein_kinase_C-phosphorylation_site
FT	Misc-difference	1386..1386	/label- protein_kinase_C-phosphorylation_site
FT	Misc-difference	1443..1443	/label- protein_kinase_C-phosphorylation_site
FT	Misc-difference	1454..1454	/label- protein_kinase_C-phosphorylation_site
FT	Misc-difference	422..422	/label- protein_kinase_C-phosphorylation_site
FT	Misc-difference	640..640	/label- protein_kinase_A-phosphorylation_site
FT	Misc-difference	685..685	/label- protein_kinase_A-phosphorylation_site
FT	Misc-difference	699..699	/label- protein_kinase_A-phosphorylation_site
FT	Misc-difference	711..711	/label- protein_kinase_A-phosphorylation_site
FT	Misc-difference	736..736	/label- protein_kinase_A-phosphorylation_site
FT	Misc-difference	767..767	/label- protein_kinase_A-phosphorylation_site
FT	Misc-difference	787..787	/label- protein_kinase_A-phosphorylation_site
FT	Misc-difference	796..796	/label- protein_kinase_A-phosphorylation_site
FT	Misc-difference	812..812	/label- protein_kinase_A-phosphorylation_site
FT	Misc-difference		/label- protein_kinase_A-phosphorylation_site
PN	W09102796-A.		
PD	07-MAR-1991.		
PF	20-AUG-1990;	90WO-CA00267.	
RR	31-AUG-1989;	89US-0401609.	
RR	22-AUG-1989;	89US-036894.	

PR	24-AUG-1989;	89US-0399945.
XX		
PA	(HSCR-) HSC RES DEV CORP.	
PA	(UNMI) UNIV OF MICHIGAN.	
XX		
XX		
PI	Tsui LC, Riordan JR, Collins FS, Rommens JM, Iannuzzi MC;	
PI	Kerem BS, Drummler ML, Buchwald M;	
XX		
DR	WPI: 1991-087280/12.	
DR	N-PSDB: AAQ11371.	
XX		
PT	Cystic fibrosis gene - used to produce prods. for screening,	
PT	detection, diagnosis, therapy and studying cystic fibrosis	
XX		
PS	Disclosure: Fig 1; 163pp. English.	
XX		
CC	This sequence lacks amino acid Phe 508 of the normal protein, as a	
CC	result of a 3bp deletion in the nucleotide sequence.	
CC	The CF gene and its gene prod., nucleic acid probes and antibodies	
CC	to the gene prod. can be used for screening and detection of CF	
CC	carriers, CF diagnosis, prenatal CF screening and diagnosis,	
CC	and gene and drug therapy. The prods. can also be used to develop	
CC	improved methods of treatment and to study the disease.	
CC	See AAQ11046 for the normal CF gene and AAQ11047-48 for CF probes.	
XX		
SO	Sequence 1479 AA;	
Query Match	94.8%; Score 110; DB 12;	
Best Local Similarity	95.5%; Pred. No. 1.1e-06;	
Matches 21; Conservative 0; Mismatches 1; Indels 0; Gaps 0.		
OY	1 GLEISEETINEDLKECFDME 22	
Db	816 GLEISEETINEDLKECFDME 837	
RESULT 9		
ARI3231		
ID	ARI3231 standard; Protein; 1479 AA.	
XX		
XX	ARI3231;	
AC		
XX		
DT	14-OCT-1991 (first entry)	
XX		
DE	CFTR delta I507.	
XX		
KW	Deletion; mutant; diagnosis; antibodies; drug therapy;	
KW	ATP-binding domain.	
XX		
OS	Homo sapiens.	
XX		
Key	Location/Qualifiers	
FT	81..102	
FT	/label= membrane-spanning_domain	
FT	118..138	
FT	/label= membrane-spanning_domain	
FT	195..215	
FT	/label= membrane-spanning_domain	
FT	221..241	
FT	/label= membrane-spanning_domain	
FT	308..328	
FT	/label= membrane-spanning_domain	
FT	330..350	
FT	/label= membrane-spanning_domain	
FT	859..879	
FT	/label= membrane-spanning_domain	
FT	911..931	
FT	/label= membrane-spanning_domain	
FT	990..1010	
FT	/label= membrane-spanning_domain	
FT	1013..1033	
FT	/label= membrane-spanning_domain	
FT	1102..1122	
FT	/label= membrane-spanning_domain	
Domain		

FT	Domain	/label= membrane-spanning_domain 1128..1149
FT	Domain	/label= membrane-spanning_domain 433..583
FT	Domain	/label= ATP-binding_domain 1218..1385
FT	Modified-site	/label= ATP-binding_domain 50
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FT	Modified-site	/label= phosphorylation_site 164
FT		/label= phosphorylation_site /note= "by protein kinases C"
FT	Modified-site	/label= phosphorylation_site 168
FT		/label= phosphorylation_site /note= "by protein kinases C"
FT	Modified-site	/label= phosphorylation_site 256
FT		/label= phosphorylation_site /note= "by protein kinases C"
FT	Modified-site	/label= phosphorylation_site 271
FT		/label= phosphorylation_site /note= "by protein kinases C"
FT	Modified-site	/label= phosphorylation_site 296
FT		/label= phosphorylation_site /note= "by protein kinases C"
FT	Modified-site	/label= phosphorylation_site 422
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FT	Modified-site	/label= phosphorylation_site 485
FT		/label= phosphorylation_site /note= "by protein kinases C"
FT	Modified-site	/label= phosphorylation_site 501
FT		/label= phosphorylation_site /note= "by protein kinases C"
FT	Modified-site	/label= phosphorylation_site 581
FT		/label= phosphorylation_site /note= "by protein kinases C"
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FT		/label= phosphorylation_site /note= "by protein kinases C"
FT	Modified-site	/label= phosphorylation_site 640
FT		/label= phosphorylation_site /note= "by protein kinases C"
FT	Modified-site	/label= phosphorylation_site 681
FT		/label= phosphorylation_site /note= "by protein kinases C"
FT	Modified-site	/label= phosphorylation_site 685
FT		/label= phosphorylation_site /note= "by protein kinases C"
FT	Modified-site	/label= phosphorylation_site 699
FT		/label= phosphorylation_site /note= "by protein kinases C and A"
FT	Modified-site	/label= phosphorylation_site 706
FT		/label= phosphorylation_site /note= "by protein kinases A"
FT	Modified-site	/label= phosphorylation_site 711
FT		/label= phosphorylation_site /note= "by protein kinases C"
FT	Modified-site	/label= phosphorylation_site 736
FT		/label= phosphorylation_site /note= "by protein kinases A"
FT	Modified-site	/label= phosphorylation_site 767
FT		/label= phosphorylation_site /note= "by protein kinases A"
FT	Modified-site	/label= phosphorylation_site 787
FT		/label= phosphorylation_site /note= "by protein kinases C"
FT	Modified-site	/label= phosphorylation_site 789
FT		/label= phosphorylation_site /note= "by protein kinases A"
FT	Modified-site	/label= phosphorylation_site 799
FT		/label= phosphorylation_site /note= "by protein kinases C"

FT	Modified-site	790	/label= phosphorylation_site
FT		/note= "by protein kinases C"	
FT	Modified-site	794	/label= phosphorylation_site
FT		/note= "by protein kinases A"	
FT	Modified-site	808	/label= phosphorylation_site
FT		/note= "by protein kinases C"	
FT	Modified-site	812	/label= phosphorylation_site
FT		/note= "by protein kinases A"	
FT	Modified-site	965	/label= phosphorylation_site
FT		/note= "by protein kinases C"	
FT	Modified-site	1057	/label= phosphorylation_site
FT		/note= "by protein kinases C"	
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FT		/note= "by protein kinases C"	
FT	Modified-site	1386	/label= phosphorylation_site
FT		/note= "by protein kinases C"	
FT	Modified-site	1443	/label= phosphorylation_site
FT		/note= "by protein kinases C"	
FT	Modified-site	1454	/label= phosphorylation_site
FT		/note= "by protein kinases C"	
PN	WO9110734-A.		
PD	25-JUL-1991.		
PF	11-JAN-1991;	91WO-CA00009.	
PR	10-JUL-1990;	90CA-2020817.	
PR	12-JAN-1990;	90CA-2007699.	
PR	01-MAR-1990;	90CA-2011253.	
PA	(HSCR-) HSC RES DEV CORP.		
PI	Tsui LC, Rommens JW, Kerem B;		
PI	WPI: 1991-238022/32.		
PI	N-PSDB; AA013053.		
PT	Mutant cystic fibrosis trans-membrane conductance regulator gene		
PT	used for producing prods. for diagnosis, screening and therapy		
PT	of cystic fibrosis		
PS	Claim 20; Page 124; 178pp; English.		
CC	The deletion of the amino acid at the 1506 or 1507 position		

CC results in the loss of an isoleucine residue from the putative
 CC CFTR, within the same ATP-binding domain where deltaF508 resides,
 CC but it is not evident whether this deleted amino acid corresponds
 CC to the position 506 or 507. Since the 506 and 507 positions are
 CC repeats, it is at present impossible to determine in which position
 CC the deletion occurs.
 CC The mutant CF gene when expressed in cells of the human body, is
 CC associated with altered cell function which correlates with the
 CC genetic disease cystic fibrosis.
 CC See also AA013053-72.

SO Sequence 1479 AA;

Query Match 94.8%; Score 110; DB 12; Length 1479;
 Best Local Similarity 95.5%; Pred. No. 1.1e-06;
 Matches 21; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

OY 1 GLEISEINEEDLKECFDDME 22
 |||||
 DB 816 GLEISEINEEDLKECLFDDME 837

RESULT 10

ID AA02279 standard; Protein; 1479 AA.

AA02279;

08-JUL-1999 (first entry)

DeltaF508 cystic fibrosis transmembrane conductance regulator.

KW Flavone; isoflavone; resveratrol; ascorbic acid; ascorbate salt;
 KW dehydroascorbic acid; chloride transport; epithelial cell;
 KW cystic fibrosis; chloride ion conductance;
 KW cystic fibrosis transmembrane conductance regulator; CFTR;
 KW chronic bronchitis; asthma; intestinal constipation.

OS Homo sapiens.

MO9918953-A1.

22-APR-1999.

16-OCT-1998; 98WO-US21887.

16-OCT-1997; 97US-0951912.

(CHIL-) CHILDREN'S HOSPITAL OAKLAND RES INST.

Fischer HB, Illek B;

WPI: 1999-277427/23.

N-PSDB; AAX35553.

Use of flavones and isoflavones - for stimulating chloride transport
 in epithelial cells and treating cystic fibrosis

Disclosure; Page 73-80; 97pp; English.

CC The specification describes compounds comprising flavones/isoflavones,
 CC resveratrol, ascorbic acid, ascorbate salts and/or dehydroascorbic
 CC acid which can be used for stimulating chloride transport in epithelial
 CC cells and treating cystic fibrosis. The compounds can be used to
 CC increase chloride ion conductance in airway epithelial cells or
 CC intestinal, pancreas, gallbladder, sweat duct, salivary gland or mammary
 CC epithelial cells. The compounds are useful for treating a patient with
 CC cystic fibrosis, where the patient's cystic fibrosis transmembrane
 CC conductance regulator (CFTR) protein has a deletion at position 508 or
 CC point mutation at 551. They may also be used for treating chronic
 CC bronchitis, asthma and intestinal constipation. The present sequence
 CC represents a human CFTR protein with a F508 deletion mutation.

SO Sequence 1479 AA;

Query Match 94.8%; Score 110; DB 20; Length 1479;
 Best Local Similarity 95.5%; Pred. No. 1.1e-06;
 Matches 21; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

OY 1 GLEISEINEEDLKECFDDME 22
 |||||
 DB 816 GLEISEINEEDLKECLFDDME 837

RESULT 11

ID AA074516 standard; Protein; 1479 AA.

AA074516;

23-APR-2002 (first entry)

Human delta-F508-CFTR mutant polypeptide.

KW Human; cystic fibrosis transmembrane conductance regulator; CFTR;
 KW Flavone; isoflavone; chloride transport; epithelial tissue; mucus;
 KW cystic fibrosis; chronic bronchitis; asthma; delta-F508-CFTR; protein.

OS Homo sapiens.

US6329422-B1.

11-DEC-2001.

16-OCT-1998; 98US-0174077.

16-OCT-1997; 97US-0951912.

(CHIL-) CHILDREN'S HOSPITAL OAKLAND RES INST.

Fischer H, Illek B;

WPI: 2002-105224/14.

N-PSDB; AAS20529.

Pharmaceutical composition for the treatment of cystic fibrosis
 comprises flavones or isoflavones

Disclosure; Column 37-44; 50pp; English.

CC The invention relates to a pharmaceutical composition comprising one or
 CC more compounds such as flavones or isoflavones, capable of stimulating
 CC chloride transport in epithelial tissues, for treatment of cystic
 CC fibrosis and other diseases associated with excessive accumulation of
 CC mucus, e.g. chronic bronchitis and asthma. The active compound increases
 CC expression of a cystic fibrosis transmembrane conductance regulator
 CC (CFTR) in an epithelial cell and/or acts as a chemical chaperone that
 CC increases trafficking of a CFTR to a plasma membrane in an epithelial
 CC cell. This sequence represents the human delta-F508-CFTR mutant
 CC polypeptide of the invention.

SO Sequence 1479 AA;

Query Match 94.8%; Score 110; DB 23; Length 1479;
 Best Local Similarity 95.5%; Pred. No. 1.1e-06;
 Matches 21; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

OY 1 GLEISEINEEDLKECFDDME 22
 |||||
 DB 816 GLEISEINEEDLKECLFDDME 837

RESULT 12

ID AAR11115 standard; Protein; 1480 AA.

AC AAR1115;
 XX 22-MAY-1991 (first entry)
 DE Cystic fibrosis transmembrane conductance regulatory protein.
 XX Cystic fibrosis; transmembrane conductance regulatory protein; CFTR;
 KM diagnosis.
 XX Homo sapiens.
 OS
 FH Key
 FH Location/Qualifiers
 FT 81..102
 FT /label= potential_membrane-spanning_segment
 FT 118..138
 FT /label= potential_membrane-spanning_segment
 FT 195..215
 FT /label= potential_membrane-spanning_segment
 FT 221..241
 FT /label= potential_membrane-spanning_segment
 FT 308..328
 FT /label= potential_membrane-spanning_segment
 FT 330..350
 FT /label= potential_membrane-spanning_segment
 FT 860..880
 FT /label= potential_membrane-spanning_segment
 FT 912..932
 FT /label= potential_membrane-spanning_segment
 FT 991..1011
 FT /label= potential_membrane-spanning_segment
 FT 1014..1034
 FT /label= potential_membrane-spanning_segment
 FT 1103..1123
 FT /label= potential_membrane-spanning_segment
 FT 1129..1150
 FT /label= potential_membrane-spanning_segment
 FT 433..584
 FT /label= putative_ATP-binding_folds
 FT 1219..1386
 FT /label= putative_ATP-binding_folds
 FT 50..50
 FT /label= protein_kinase_C-phosphorylation_site
 FT 63..63
 FT /label= protein_kinase_C-phosphorylation_site
 FT 164..164
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 FT 168..168
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 FT 256..256
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 FT 271..271
 FT /label= protein_kinase_C-phosphorylation_site
 FT 296..296
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 FT 485..485
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 FT 501..501
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 FT 582..582
 FT /label= protein_kinase_C-phosphorylation_site
 FT 623..623
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 FT 682..682
 FT /label= protein_kinase_C-phosphorylation_site
 FT 686..686
 FT /label= protein_kinase_C-phosphorylation_site
 FT 707..707
 FT /label= protein_kinase_C-phosphorylation_site
 FT 790..790
 FT /label= protein_kinase_C-phosphorylation_site
 FT 791..791
 FT /label= protein_kinase_C-phosphorylation_site
 FT 809..809
 FT Misc-difference

FT /label= protein_kinase_C-phosphorylation_site
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 FT 1178..1178
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 FT 1248..1248
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 FT 1387..1387
 FT /label= protein_kinase_C-phosphorylation_site
 FT 1444..1444
 FT /label= protein_kinase_C-phosphorylation_site
 FT 1455..1455
 FT /label= protein_kinase_C-phosphorylation_site
 FT 422..422
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 FT 641..641
 FT /label= protein_kinase_A-phosphorylation_site
 FT 686..686
 FT /label= protein_kinase_A-phosphorylation_site
 FT 700..700
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 FT 712..712
 FT /label= protein_kinase_A-phosphorylation_site
 FT 737..737
 FT /label= protein_kinase_A-phosphorylation_site
 FT 768..768
 FT /label= protein_kinase_A-phosphorylation_site
 FT 788..788
 FT /label= protein_kinase_A-phosphorylation_site
 FT 795..795
 FT /label= protein_kinase_A-phosphorylation_site
 FT 813..813
 FT /label= protein_kinase_A-phosphorylation_site
 FT 508..508
 FT /label= mutation
 FT /note= "residue is deleted in the CFTR mutant"
 PN WO9102796-A.
 PD 07-MAR-1991.
 PD 20-AUG-1990; 90WO-CA00267.
 PF 31-AUG-1989; 89US-0401609.
 PR 22-AUG-1989; 89US-0396894.
 PR 24-AUG-1989; 89US-0399945.
 PA (HSCR-) HSC RES DEV CORP.
 PA (UNMI) UNIV OF MICHIGAN.
 XX Tsui LC, Riordan JR, Collins FS, Rommens JM, Tannuzzi MC;
 PI Kerem BS, Drumml ML, Buchwald M;
 XX WPI; 1991-087280/12.
 DR P-PSDB; AAR1115.
 XX Cystic fibrosis gene - used to produce prods. for screening,
 PT detection, diagnosis, therapy and studying cystic fibrosis
 XX Disclosure; Fig 1, 163pp; English.

CC The protein has a mol.wt. of 170 kD and has two repeated motifs,
 CC each comprising a set of amino acids comprising six highly
 CC hydrophobic segments (see membrane spanning segments in features),
 CC capable of spanning a lipid bilayer of an epithelial cell membrane
 CC several times followed by an amino acid sequence constituting
 CC a nucleotide (ATP)-binding fold. (The first of the (ATP)-binding
 CC folds has the p1e deletion (See AAR11602 for the mutant protein)).
 CC Between the repeated motifs is a highly charged cytoplasmic domain.
 CC The CF gene and its gene prod., nucleic acid probes and antibodies
 CC to the gene prod. can be used for screening and detection of CF
 CC carriers, CF diagnosis, prenatal CF screening and diagnosis of CF
 CC and gene and drug therapy. The prods. can also be used to develop
 CC improved methods of treatment and to study the disease.
 CC See AAQ11371 for the mutant CF gene and AAQ11047-48 for CF probes.

CC Sequence 1480 AA;

Query Match 94.88; Score 110; DB 12; Length 1480;
 Best Local Similarly 95.58; Pred. No. 1.1e-06;
 Matches 21; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 GLEISERINEDLKECFDME 22
 |||||

Db 817 GLEISERINEDLKECLFDME 838

RESULT 13

AAAR13232 ID AAR13232 standard; Protein; 1480 AA.

XX AAR13232;

XX 14-OCT-1991 (first entry)

DE CFTR G85E.

KW Deletion; mutant; diagnosis; antibodies; drug therapy.

XX Homo sapiens.

XX Key Location/Qualifiers

FT Domain /label= membrane-spanning_domain
 FT 81..102
 FT Domain /label= membrane-spanning_domain
 FT 118..138
 FT Domain /label= membrane-spanning_domain
 FT 195..215
 FT Domain /label= membrane-spanning_domain
 FT 221..241
 FT Domain /label= membrane-spanning_domain
 FT 308..328
 FT Domain /label= membrane-spanning_domain
 FT 330..350
 FT Domain /label= membrane-spanning_domain
 FT 860..880
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 FT 912..932
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 FT 991..1011
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 FT 1014..1034
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 FT 1103..1123
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 FT 1129..1150
 FT Domain /label= membrane-spanning_domain
 FT 433..584
 FT Domain /label= ATP-binding_domain
 FT 1219..1386
 FT Modified-site /label= ATP-binding_domain
 FT 50
 FT Modified-site /label= phosphorylation_site
 FT /note= "by protein kinases C"
 FT 63
 FT Modified-site

FT /label= phosphorylation_site
 FT /note= "by protein kinases C"
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 FT Modified-site /label= phosphorylation_site
 FT /note= "by protein kinases C"
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 FT 813
 FT Modified-site /label= phosphorylation_site


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XX      WO9110734-A.
PN
XX      25-JUL-1991.
PD
XX      11-JAN-1991; 91WO-CA00009.
PE
XX      10-JUL-1990; 90CA-2020817.
PR      12-JAN-1990; 90CA-2007699.
PR      01-MAR-1990; 90CA-2011253.
XX
PA      (HSCR-) HSC RES DEV CORP.
XX
PI      Tsui LC, Rommens JM, Kerem B;
DR      MPI; 1991-238022/32.
DR      N-PSDB; AAQ13056.
XX
PT      Mutant cystic fibrosis trans-membrane conductance regulator gene
PT      used for producing prods. for diagnosis, screening and therapy
XX      of cystic fibrosis
XX
PS      Claim 20; Page 124; 178pp; English.
XX
CC      The G178R mutation in exon 5 involves a G to A transition at
CC      nucleotide position 664 resulting in a Gly to Arg change at amino
CC      acid position 178.
CC      The mutant CF gene when expressed in cells of the human body, is
CC      associated with altered cell function which correlates with the
CC      genetic disease cystic fibrosis.
CC      See also AAQ13053-72.
XX
SQ      Sequence 1480 AA;

```

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Query Match          94.8%; Score 110; DB 12; length 1480;
Best Local Similarity 95.5%; Pred. No. 1,1e-06;
Matches 21; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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QY      1 GLEISEINEDLKECFDDME 22
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Db      817 GLEISEINEDLKECFDDME 838

```

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Search completed: July 16, 2003, 14:01:12
Job time : 70 secs

```

